Appln # 09/765,865
Paper # 12 Attach.

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FILE 'MEDLINE, CAPLUS, BIOSIS, EMBASE' ENTERED AT 18:50:38 ON 22 SEP 2002

L1 321 P15A
L2 30 (HOST RANGE OR HOST-RANGE) AND L1
L3 9 DUP REM L2 (21 DUPLICATES REMOVED)
L4 0 RK2 REPLICON AND GLUCOSE DEHYDROGENASE
L5 98 (RK2 OR RK4) (A) (REPLICON OR ORI)
L6 0 L5 AND "GLUCOSE DEHYDROGENASE"
L7 0 L5 AND DEHYDROGENASE
L8 147 BROAD-HOST-RANGE AND DEHYDROGENASE
L9 67 DUP REM L8 (80 DUPLICATES REMOVED)
L10 74 (BROAD-HOST-RANGE (S) VECTOR) (P) DEHYDROGENASE
L11 25 DUP REM L10 (49 DUPLICATES REMOVED)
L12 0 L11 AND (NON-MOBILIZABLE OR NONMOBILIZABLE OR NON-MOBILISABLE O

**Loeb, Bronw n**

**From:** Loeb, Bronwen
**Sent:** Sunday, September 22, 2002 7:44 PM
**To:** STIC-ILL
**Subject:** ILL order 09/765,8965

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Appln 09/765,865

Sykes et al (1988) Methods Enzymol. (1988), 166(Branched Chain Amino Acids), 350-9 BCKAD
Sykes et al (1987) JOURNAL OF BACTERIOLOGY, (1987 Apr) 169 (4) 1619-25. branched chain keto acid dehydrog. (=BCKAD)
Lehrbach et al (1984) JOURNAL OF BACTERIOLOGY, (1984 Jun) 158 (3) 1025-32.
Bagadasarian et al (1981) Gene 16:237-247 (pKT230 and pKT210 construction) probably $Mob^+$

dihydrodihydroxybenzoic acid dehydrog.

Deretic et al (1987) JOURNAL OF BACTERIOLOGY, (1987 Jan) 169 (1) 351-8. GDPmannose dehydrogenase
Bagdasarian et al., Gene (1983)26:273-282 (pMMb24 construction)

Deretic et al (1987) GENE, (1987) 57 (1) 61-72. GDPmannose dehydrogenase